

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: US1091765,555B

Source: ITFW16

Date Processed by STIC: 12-16-09

ENTERED



IFW16

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/765,555B

DATE: 12/16/2004

TIME: 10:54:13

Input Set : D:\27801-20014.20 - Sub Seqlist.txt
 Output Set: N:\CRF4\12162004\I765555B.raw

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3 <110> APPLICANT: BARBAS, Carlos
4      STEGE, Justin
5      GUAN, Xueni
6      DALMIA, Bipin
8 <120> TITLE OF INVENTION: METHODS AND COMPOSITIONS TO MODULATE
9      EXPRESSION IN PLANTS
11 <130> FILE REFERENCE: 27801-20014.20
13 <140> CURRENT APPLICATION NUMBER: 09/765,555B
14 <141> CURRENT FILING DATE: 2001-01-19
16 <150> PRIOR APPLICATION NUMBER: 09/620,897
17 <151> PRIOR FILING DATE: 2000-07-21
19 <150> PRIOR APPLICATION NUMBER: US 60/177,468
20 <151> PRIOR FILING DATE: 2000-01-21
22 <160> NUMBER OF SEQ ID NOS: 78
24 <170> SOFTWARE: FastSEQ for Windows Version 4.0
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 532
28 <212> TYPE: DNA
29 <213> ORGANISM: Artificial Sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: Promoter CsvMV
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36 acggaaaaaa ctatgaaagt attatgttag ctcagcaaga agcagatcaa tatgcggcac      120
37 atatgcaacc tatgttcaaa aatgaagaat gtacagatac aagatcctat actgccagaa      180
38 tacgaagaag aatacgtaga aattgaaaaaa gaagaaccag gcgaagaaaaa gaatcttcaa      240
39 gacgtaagca ctgacgacaa caatgaaaaag aagaagataa ggtcggttat tgtgaaagag      300
40 acatagagga cacatgtaag gtggaaaatg taaggcgga aagtaacctt atcacaaagg      360
41 aatcttatcc cccactactt atccttttat attttccgt gtcatttttgc cccttgagtt      420
42 ttccatatata aggaaccaag ttccgcattt gtgaaaaacaa gaaaaaattt ggtgtaagct      480
43 atttcttttg aagtactgag gataacaattt cagagaaattt tgtaagtttta      532
45 <210> SEQ ID NO: 2
46 <211> LENGTH: 18
47 <212> TYPE: DNA
48 <213> ORGANISM: Artificial Sequence
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Zinc finger protein 2C7 binding site
53 <400> SEQUENCE: 2
54 gcgtggcgaa cgtggcg      18
56 <210> SEQ ID NO: 3
57 <211> LENGTH: 51
58 <212> TYPE: DNA
59 <213> ORGANISM: Artificial Sequence

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114	aaggccctatg	cttgcctgt	cgagtcctgc	gatcgccgct	tttctaagtc	ggctgatctg	2340
115	aaggcgcata	tccgcatacca	cacaggccag	aaggcctcc	agtgtcaat	atgcatgcgt	2400
116	aacttcagtc	gtatgtacca	ccttaccacc	cacatccgca	cccacacagg	cgagaagcct	2460
117	tttgcctgtg	acatttgtgg	gaggaagtt	gccaggagtg	atgaacgcaa	gaggcatacc	2520
118	aaaatccatt	taagacagaa	ggactctaga	actagtggcc	aggccgcca	ggctagcccc	2580
119	aaaaagaaaac	gcaaagttgg	gcgcgcccac	gcgcgtggacg	atttcgatct	cgacatgctg	2640
120	ggttctgtatg	ccctcgatga	ctttgacctg	gatatgttg	gaagcgacgc	atggatgac	2700
121	tttgcatacg	acatgtcg	ctccgcatact	ctggacgatt	tcgatctcg	tatgttaatt	2760
122	aactaccgt	acgacgttcc	ggactacgt	tcttgcataat	tcgcggccgc	ggggccgagc	2820
123	ctagggagga	gctcaagatc	ccccgaaattt	ccccgatcgt	tcaaacaattt	ggcaataaaag	2880
124	tttcttaaga	ttgaatcctg	ttgcgggtct	tgcgatgatt	atcatctaatt	ttctgttgaa	2940
125	ttacgttaag	catgtataaa	ttaacatgtat	atgcatgacg	ttatttatga	gatgggtttt	3000
126	tatgattaga	gtcccgcaat	tatacatatta	atacgcgata	gaaaacaaaa	tatagcgcgc	3060
127	aaacttagat	aaattatcgc	gcgcgggtgc	atctatgtta	ctagatccgg	gaattggta	3120
128	C						3121
130	<210>	SEQ ID NO:	5				
131	<211>	LENGTH:	3069				
132	<212>	TYPE:	DNA				
133	<213>	ORGANISM:	Artificial Sequence				
135	<220>	FEATURE:					
136	<223>	OTHER INFORMATION:	pND3018 coding redion				
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140	aaattaccac	atatttttt	tgtcacactt	gttgaagt	cagtttatct	atctttatac	120
141	atatatattaa	actttaact	acgaaataata	taatctatag	tactacaata	atatcgtgt	180
142	tttagagaat	catataaatg	aacagttaga	catggctaa	aggacaattt	agtattttga	240
143	caacaggact	ctacagtttt	atctttttag	tgtgcatagt	ttctcctttt	tttttgc当地	300
144	tagttcacc	tatataatac	ttcatccatt	ttatttagtac	atccatttag	ggttttaggt	360
145	taatggttt	tatagactaa	tttttttagt	acatctattt	tattctattt	tagcctctaa	420
146	attaagaaaa	ctaaaactct	attttagttt	tttttattaa	taatttagat	ataaaataga	480
147	ataaaataaa	gtgactaaaa	attaaacaaa	taccctttaa	gaaattaaaa	aaactaagga	540
148	aacattttc	ttgttcgag	tagataatgc	cagcctgtta	aacgcgcgtc	acgagtctaa	600
149	cgAACACCAA	ccagcgaacc	agcagcgtc	cgtcgggcca	agcgaagcag	acggcacggc	660
150	atctctgtcg	ctgcctctgg	acccctctcg	agagttccgc	tccaccgtt	gacttgctcc	720
151	gctgtcgcca	tccagaaatt	gcgtggcgga	gcggcagacg	tgagccggca	cggcaggcgg	780
152	cctccctcc	ctctcacggc	acggcagacta	cgggggattt	cttcccacc	gctcttcgc	840
153	tttccctcc	tcgcccggcg	taataaataag	acacccccctc	cacacccctt	ttcccccaacc	900
154	tcgtgttgtt	cgagcgcac	acacacacaa	ccagatctcc	cccaaatttca	cccgctggca	960
155	cctccgttcc	aaggtacgcc	gctcgcttc	cccccccccc	cctctctacc	ttctctatgt	1020
156	cgccgttccg	gtccatggtt	aggggccgg	agtttactt	ctgttcatgt	ttgtgtttaga	1080
157	tccgtgtttt	tgtagatcc	gtgctgtat	cggtcgatca	cggtatgcgc	ctgtacgtca	1140
158	gacacgttct	gattgctaac	ttgcgcgtgt	ttctctttgg	ggaatcttgg	gatggctcta	1200
159	gccgttccgc	agacgggatc	gatttcatga	ttttttttgt	ttcggttgc	agggtttgg	1260
160	ttgcctttt	ccttatttc	aatatatgtcc	gtgcacttgt	ttgtcggttgc	atcttttcat	1320
161	gtttttttt	gtcttgggtt	tgatgtatgt	gtcttgggtt	gcggcggtt	tagatcgag	1380
162	tagaattctg	tttcaaacta	cctgggtggat	ttattaattt	tggatctgtat	tgtgtgtgcc	1440
163	atacatattc	atagttacga	attgaagatg	atggatggaa	atatcgatct	aggataggt	1500
164	tacatgttga	tgcggttttt	actgtatgtat	atacagagat	gttttttgtt	cgcttgggttgc	1560
165	tgtatgtatgt	gtgtgggttgg	gcggcggttgc	attcgatctat	gatcgagat	gaataactgtt	1620

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166	tcaaactacc	tggtgttattt	attaattttg	gaactgtatg	tgtgtgtcat	acatcttc	cat	1680							
167	agttacgagt	ttaagatgga	tggaaatatac	gatctaggat	aggatacat	gtt	gtgtgg	1740							
168	gttttactga	tgcata	tataca	tgtggcata	tgcagcatct	attc	atata	1800							
169	agtacccatc	tattataata	aacaagtatg	tttataatt	atttt	gatct	tgtatata	1860							
170	gatgtatggc	atatgcagca	gctatatgtg	gat	tttttta	gc	cctt	catacgctat	1920						
171	ttat	ttgc	tgtt	ctttgtcg	tg	ctcac	ct	gtt	tttggt	gt	tacttctg	1980			
172	cagg	tcgact	ctagg	actagtgag	ccatgggcta	gcat	gggc	gc	tgccgtgc	gc	2040				
173	atgaacatcc	agatgtgt	cgaagccgt	gattatctgg	aac	gcggg	ga	gc	gcgaagcc	2100					
174	gagcacggct	acgccc	acat	gctgccat	ccgaaa	aac	gcaaggt	gg	cccaggcg	2160					
175	gc	cctcg	gagc	tccctatgc	ttgc	cc	cctgc	g	atcgc	ctt	ctaa	2220			
176	gt	gtatctga	agc	ccat	ccac	ac	aggccaga	ag	cccttcca	gt	gtcgaata	2280			
177	tgc	atgc	gt	actt	cagtc	tagt	gaccac	ct	accac	cc	acacaggc	2340			
178	gaga	gccc	tt	tgctgt	ga	cattt	gtgg	ag	gaagttt	cc	aggagt	ta	aacgc	2400	
179	aggc	cata	cca	aa	atccatac	cgg	tgaga	cc	ctatg	ct	cc	gtc	gtc	2460	
180	cgcc	gctt	ttt	cta	agtc	cg	g	tc	at	ccat	at	cc	gc	aggccagaag	2520
181	cc	c	tttcc	ca	gt	cg	at	at	at	at	cc	at	cc	acc	2580
182	atcc	gcac	cc	ac	acagg	gc	aa	gc	ctt	gt	gt	gt	gac	tttggag	2640
183	agg	gtat	g	a	ac	ca	ag	g	at	ccat	tt	aa	gac	aga	2700
184	agt	ggcc	cagg	cc	ggc	at	cc	g	act	tc	tt	tt	tt	gg	2760
185	acc	gag	ctcg	g	at	cccc	cg	at	tt	cc	cc	at	tt	tt	2820
186	aag	att	gaat	c	c	tgtt	cc	gt	tt	tc	at	at	tc	tgt	2880
187	taa	g	gat	g	taa	taa	ca	ta	at	at	at	at	at	at	2940
188	tag	at	cc	cg	ca	att	tata	ta	taa	at	ac	aa	at	at	3000
189	ggata	aaat	at	tc	gc	gc	gc	gg	tgt	cat	ct	at	at	at	3060
190	cc	agc	gg	gc	cc	gg	gg	gg	t	act	at	at	at	at	3069
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193	<211>	LENGTH:	156												
194	<212>	TYPE:	DNA												
195	<213>	ORGANISM:	Artificial Sequence												
197	<220>	FEATURE:													
198	<223>	OTHER INFORMATION:	6X2C7 binding site												
200	<400>	SEQUENCE:	6												
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202	gc	gg	gg	gt	gg	gg	gg	gg	gg	gg	gg	gg	gg	gg	120
203	tgg	gg	ca	ac	ac	gg	gg	gg	gg	gg	gg	gg	gg	gg	156
205	<210>	SEQ ID NO:	7												
206	<211>	LENGTH:	18												
207	<212>	TYPE:	DNA												
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210	<220>	FEATURE:													
211	<223>	OTHER INFORMATION:	ZFPAp3												
213	<400>	SEQUENCE:	7												
214	gat	gg	ag	tt	g	a	aa	aa	gt	a	aa	aa	gt	a	18
216	<210>	SEQ ID NO:	8												
217	<211>	LENGTH:	21												
218	<212>	TYPE:	DNA												
219	<213>	ORGANISM:	Artificial Sequence												
221	<220>	FEATURE:													
222	<223>	OTHER INFORMATION:	ZFP from -85 to -65												

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224 <400> SEQUENCE: 8
225 gcctccttcc tcctctca c
227 <210> SEQ ID NO: 9
228 <211> LENGTH: 18
229 <212> TYPE: DNA
230 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: ZFPm1 from -68 to -85
235 <400> SEQUENCE: 9
236 tgagaggagg aaggaggc
238 <210> SEQ ID NO: 10
239 <211> LENGTH: 18
240 <212> TYPE: DNA
241 <213> ORGANISM: Artificial Sequence
243 <220> FEATURE:
244 <223> OTHER INFORMATION: ZFPm2 from -65 to -82
246 <400> SEQUENCE: 10
247 gagtgagagg aggaagga
249 <210> SEQ ID NO: 11
250 <211> LENGTH: 24
251 <212> TYPE: DNA
252 <213> ORGANISM: Artificial Sequence
254 <220> FEATURE:
255 <223> OTHER INFORMATION: ZFP from 294 to 317
257 <400> SEQUENCE: 11
258 gccaaactact acggctccct cacc
260 <210> SEQ ID NO: 12
261 <211> LENGTH: 18
262 <212> TYPE: DNA
263 <213> ORGANISM: Artificial Sequence
265 <220> FEATURE:
266 <223> OTHER INFORMATION: ZFPm3 from 311 to 294
268 <400> SEQUENCE: 12
269 ggagccgtag tagttggc
271 <210> SEQ ID NO: 13
272 <211> LENGTH: 18
273 <212> TYPE: DNA
274 <213> ORGANISM: Artificial Sequence
276 <220> FEATURE:
277 <223> OTHER INFORMATION: ZFPm4 from 317 to 300
279 <400> SEQUENCE: 13
280 ggtgagggag ccgttagta
282 <210> SEQ ID NO: 14
283 <211> LENGTH: 3300
284 <212> TYPE: DNA
285 <213> ORGANISM: Artificial Sequence
287 <220> FEATURE:
288 <223> OTHER INFORMATION: Partial sequence of pMal-m1 and zinc finger
289 protein ZFPm1

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VERIFICATION SUMMARY

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